

FIG. 1

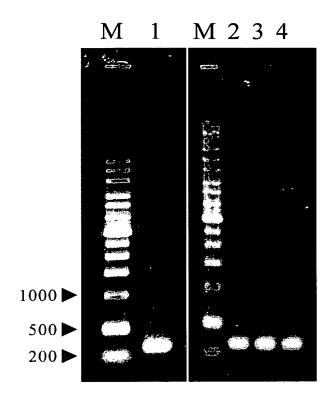


FIG. 2

		1 60
CCRC12913	(1)	
CCRC14001	(1)	
GM-080	(1)	
CCRC16100	(1)	
KLB58	(1)	
PB4	(1)	ATCCTGGCTCAGGATGAACGCTGGCGGCGTGCCTAATACATGCAAGTCGAACGAGTTCTC
F31	(1)	AACGCTGGCGGCGTGCCTAATACATGCAAGTCGAACGAGTTCTC
Consensus	(1)	
	(.)	61 120
CCRC12913	(1)	GGTGAGTAC
CCRC14001		GTGAGTAC
GM-080	(1)	GCGAGGGTGAGTAC
CCRC16100	(1)	
KLB58	(43)	
PB4	(61)	GTTGATGATCGTGCTTGCACCGAGATTCAACATGGAACGAGTGGCGGACGGGTGAGTAA GTTGATGATCGTGCTTGCACCGAGATTCAACATGGAACGAGTGGCGGACGGGTGAGTAA
F31	(45)	G GGGTGAGTA
Consensus	(61)	G GGGGAGTA
		121 180
CCRC12913	(1)	TGGGTA-CCTGCCCTTAAGTGGGGGATAACATTTGGAAACAGATGCTAATACCGCA
CCRC14001	(10)	
GM-080		CGTGGGTA-CCTGCCCTTA-GTGGGGGATAACATTTGGAAACAGATGCTAATACCGCA
CCRC16100		CGTGGGTA-CCTGCCCTTAAGTGGGGGATAACATTTGGAAACAGATGCTAATACCGCA
KLB58		CACGTGGGTAACCTGCCCTTAAGTGGGGGATAACATTTGGAAACAGATGCTAATACCGCA
PB4		CACGTGGGTAACCTGCCCTTAAGTGGGGGATAACATTTGGAAACAGATGCTAATACCGCA
F31		CACGTGGGTAACCTGCCCTTAAGTGGGGGATAACATTTGGAAACAGATGCTAATACCGCA
Consensus		C GTGGGTA CCTGCCCTTAAGTGGGGGATAACATTTGGAAACAGATGCTAATACCGCA
consensus	(101)	0.000
		181 240
CCRC12913	(56)	TAGATCCAAGAACCGCATGGTTCTTGGCTGAAAGATGGCGTAAGCTATCGCTTTTGGATG
CCRC14001	(67)	TAGATCCAAGAACCGCATGGTTCTTGGCTGAAAGATGGCGTAAGCTATCGCTTTTGGATG
GM-080	(65)	TAGATCCAAGAACCGCATGGTTCTTGGCTGAAAGATGGCGTAAGCTATCGCTTTTGGATG
CCRC16100	(72)	TAGATCCAAGAACCGCATGGTTCTTGGCTGAAAGATGGCGTAAGCTATCGCTTTTGGATG
KLB58	(163)	TAGATCCAAGAACCGCATGGTTCTTGGCTGAAAGATGGCGTAAGCTATCGCTTTTGGATG
PB4	(181)	${\tt TAGATCCAAGAACCGCATGGTTCTTGGCTGAAAGATGGCGTAAGCTATCGCTTTTGGATG}$
F31	(165)	TAGATCCAAGAACCGCATGGTTCTTGGCTGAAAGATGGCGTAAGCTATCGCTTTTGGATG
Consensus	(181)	${\tt TAGATCCAAGAACCGCATGGTTCTTGGCTGAAAGATGGCGTAAGCTATCGCTTTTGGATG}$
		241 300
CCRC12913		${\tt GACCCGCGGCGTATTAGCTAGTTGGTGAGGTAATGGCTCACCAAGGCGATGATACGTAGC}$
CCRC14001		GACCCGCGGCGTATTAGCTAGTTGGTGAGGTAATGGCTCACCAAGGCGATGATACGTAGC
GM-080		GACCCGCGGCGTATTAGCTAGTTGGTGAGGTAATGGCTCACCAAGGCGATGATACGTAGC
CCRC16100		GACCCGCGGCGTATTAGCTAGTTGGTGAGGTAATGGCTCACCAAGGCGATGATACGTAGC
KLB58		GACCCGCGGCGTATTAGCTAGTTGGTGAGGTAATGGCTCACCAAGGCGATGATACGTAGC
PB4		GACCCGCGGCGTATTAGCTAGTTGGTGAGGTAATGGCTCACCAAGGCGATGATACGTAGC
F31		GACCCGCGGCGTATTAGCTAGTTGGTGAGGTAATGGCTCACCAAGGCGATGATACGTAGC
Consensus	(241)	${\tt GACCCGCGGCGTATTAGCTAGTTGGTGAGGTAATGGCTCACCAAGGCGATGATACGTAGC}$
		200
000010010	(150)	360
CCRC12913		CGAACTGAGAGGTTGATCGGCCACATTGGGACTGAGACACGGCCCAAACTCCTACGGGAG
CCRC14001		CGAACTGAGAGGTTGATCGGCCACATTGGGACTGAGACACCGCCCAAACTCCTACGGAG
GM-080		CGAACTGAGAGGTTGATCGGCCACATTGGGACTGAGACACCGCCCAAACTCCTACGGN-
CCRC16100		CGAACTGAGAGGTTGATCGGCCACATTGGGACTGAGACACCGCCCAAACTCCTACGGG-
KLB58		CGAACTGAGAGGTTGATCGGCCACATTGGGACTGAGACACGGCCCAAACTCCTACGGG-
PB4		CGAACTGAGAGGTTGATCGGCCACATTGGGACTGAGACACGGCCCAAACTCCTACGGG-
F31		CGAACTGAGAGGTTGATCGGCCACATTGGGACTGAGACACGGCCCAAACTCCTACGGG-
Consensus	(301)	CGAACTGAGAGGTTGATCGGCCACATTGGGACTGAGACACGGCCCAAACTCCTACGGG

FIG. 3

		361 420
CCRC12913	(236)	GGCCAGCAGT-GGGAG
CCRC14001	(247)	GGGCAGCAGT-GGGAG
GM-080	(243)	GGGCAGCAGT-GGGA
CCRC16100	(250)	AGGCAGCACT-GGGA
KLB58	(341)	AGGCAGCAGTAGGGAATCTTCCACAATGGACGCAAGTCTGATGGAGCAACGCCGCGTGAG
PB4	(359)	AGGCAGCAGTAGGGAATCTTCCACAATGGACGCAAGTCTGATGGAGCAACGCCGCGTGAG
F31	(343)	AGGCAGCAGTAGGGAATCTTCCACAATGGACGCAAGTCTGATGGAGCAACGCCGCGTGAG
Consensus	(361)	AGGCAGCAGT GGGA
		421 480
CCRC12913	(251)	
CCRC14001	(262)	
GM-080	(257)	
CCRC16100	(264)	
KLB58	(401)	TGAAGAAGGCTTTCGGGTCGTAAAACTCTGTTGTTGGAGAAGAATGGTCGGCAGAGTAAC
PB4	(419)	TGAAGAAGGCTTTCGGGTCGTAAAACTCTGTTGTTGGAGAAGAATGGTCGGCAGAGTAAC
F31	(403)	TGAAGAAGGCTTTCGGGTCGTAAAACTCTGTTGTTGGAGAAGAATGGTCGGCAGAGTAAC
Consensus	(421)	
		481 540
CCRC12913	(251)	
CCRC14001	(262)	
GM-080	(257)	
CCRC16100	(264)	
KLB58	(461)	TGTTGTCGGCGTGACGGTATCCAACCAGAAAGCCACGGCTAACTACGTGCCAGCAGCCGC
PB4	(479)	TGTTGTCGGCGTGACGGTATCCAACCAGAAAGCCACGGCTAACTACGTGCCAGCAGCCGC
F31	(463)	TGTTGTCGGCGTGACGGTATCCAACCAGAAAGCCACGGCTAACTACGTGCCAGCAGCCGC
Consensus	(481)	
	(==-)	541 600
CCRC12913	(251)	
CCRC14001		
GM-080	(257)	
CCRC16100	(264)	
KLB58	(521)	GGTAATACGTAGGTGGCAAGCGTTATCCGGATTTATTGGGCGTAAAGCGAGCG
PB4	(539)	
F31	(523)	GGTAATACGTAGGTGGCAAGCGTTATCCGGATTTATTGGGCGTAAAGCGAGCG
Consensus	(541)	
		601 660
CCRC12913	(251)	
CCRC14001	(262)	
GM-080	(257)	
CCRC16100	(264)	
KLB58	(581)	TTTTTTAAGTCTGATGTGAAAGCCCTCGGCTTAACCGAAGAAGCGCATCGGAAACTGGGA
PB4	(539)	
F31	(583)	TTTTTTAAGTCTGATGTGAAAGCCCTCGGCTTAACCGAGGAAGCGCATCGGAAACTGGGA
Consensus	(601)	
		661 720
CCRC12913	(251)	661 720
CCRC12913 CCRC14001	(262)	
CCRC14001	(262)	
CCRC14001 GM-080	(262)	
CCRC14001	(262) (257) (264)	
CCRC14001 GM-080 CCRC16100	(262) (257) (264)	
CCRC14001 GM-080 CCRC16100 KLB58	(262) (257) (264) (641) (539)	
CCRC14001 GM-080 CCRC16100 KLB58 PB4	(262) (257) (264) (641) (539)	AACTTGAGTGCAGAAGAGGACAGTGGAACTCCATGTGTAGCGGTGAAATGCGTAGATATA

FIG. 3 (continued)

		721	780
CCRC12913	(251)		
CCRC14001	(262)		
GM-080	(257)		
CCRC16100	(264)		
KLB58	(701)	TGGAAGAACACCAGTGGCGAAGGCGGCTGTCTGGTCTGTAACTGACGCTGAGGCT	i'CGAAA
PB4	(539)		
F31	(703)	TGGAAGAACACCAGTGGCGAAGGCGGCTGTCTGGTCTGTAACTGACGCTGAGGCT	r CGAAA
Consensus	(721)		
		781	840
CCRC12913	(251)		
CCRC14001	(262)		
GM-080	(257)		
CCRC16100	(264)		
KLB58		GCATGGGTAGCGAACAGGATTAGATACCCTGGTAGTCCATGCCGTAAACGATGAA	ATGCTA
PB4	(539)		
F31	(763)	GCATGGGTAGCGAACAGGATTAGATACCCTGGTAGTCCATGCCGTAAACGATGAA	ATGCTA
Consensus	(781)		
		841	900
CCRC12913			
CCRC14001	(262)		
GM-080	(257)		
CCRC16100	(264)		
KLB58		GGTGTTGGAGGGTTTCCGCCCCTTCAGTGCCGCAGCTAACGCATTAAGCATTCCGC	
PB4	(539)		
F31		GGTGTTGGAGGGTTTCCGCCCTTCAGTGCCGCAGCTAACGCATTAAGCATTCCGC	JC1666
Consensus	(841)		
		901	960
CCRC12913	(251)		
CCRC14001	(262)		
GM-080	(257)		
CCRC16100	(264)		
KLB58	(881)		GTGGAG
PB4	(539)		
F31		GAGTACGACCGCAAGGTTGAAACTCAAAGGAATTGACGGGGGCCCGCACAAGCGC	GTGGAG
Consensus	(901)		
		961	1020
CCRC12913	(251)		
CCRC14001	(262)		
GM-080	(257)		
CCRC16100	(264)		
KLB58	(941)	CATGTGGTTTAATTCGAAGCAACGCGAAGAACCTTACCAAGTCTTGACATCTTTT	FGATCA
PB4	(539)		
F31	(943)	CATGTGGTTTAATTCGAAGCAACGCGAAGAACCTTACCAAGTCTTGACATCTTTT	FGATCA
Consensus	(961)		
	()	1021	1080
CCRC12913	(251)		
CCRC14001	(262)		
GM-080	(257)		
CCRC16100		CCTGAGAGATCAGGTTTCCCCTTCGGGGGCAAAATGACAGGTGGTGCATGGTTG	
KLB58			- -
PB4 F31	(539)	CCTGAGAGATCAGGTTTCCCCTTCGGGGGCAAAATGACAGGTGGTGCATGGTTC	TCGTCA
Consensus	(1003)		. 55 . 6.1
consensus	(1021)		

FIG. 3 (continued)

		1081 1140
CCRC12913	(251)	
CCRC14001	(262)	
GM-080	(257)	
CCRC16100	(264)	
KLB58	(1061)	GCTCGTGTGGGATGTTGGGTTAAGTCCCGCAACGAGCGCAACCCTTATGACTAGTTG
PB4	(539)	
F31		GCTCGTGTCGTGAGATGTTGGGTTAAGTCCCGCAACGAGCGCAACCCTTATGACTAGTTG
Consensus	(1081)	
		1141 1200
CCRC12913	(251)	
CCRC14001	(262)	
GM-080	(257)	
CCRC16100	(264)	
KLB58	(1121)	CCAGCATTTAGTTGGGCACTCTAGTAAGACTGCCGGTGACAAACCGGAGGAAGGTGGGGA
PB4	(539)	
F31	(1123)	CCAGCATTTAGTTGGGCACTCTAGTAAGACTGCCGGTGACAAACCGGAGGAAGGTGGGGA
Consensus	(1141)	
	, ,	
		1201 1260
CCRC12913		
CCRC14001	(262)	
GM-080	(257)	
CCRC16100	(264)	
KLB58	(1181)	TGACGTCAAATCATCATGCCCCTTATGACCTGGGCTACACACGTGCTACAATGGATGG
PB4	(539)	TO A CORTO A TO A TO A TO A TO A CORTO A TO A CORTO A
F31	(1183)	TGACGTCAAATCATCATGCCCCTTATGACCTGGGCTACACACGTGCTACAATGGATGG
Consensus	(1201)	
		1261 1320
CCRC12913	(251)	
CCRC14001	(262)	
GM-080	(257)	
CCRC16100	(264)	
KLB58	(1241)	${\tt CAACGAGTTGCGAGACCGCGAGGTCAAGCTAATCTCTTAAAGCCATTCTCAGTTCGGACT}$
PB4	(539)	
F31	(1243)	CAACGAGTTGCGAGACCGCGAGGTCAAGCTAATCTCTTAAAGCCATTCTCAGTTCGGACT
Consensus	(1261)	
		1200
CCDC19019	(251)	1321 1380
CCRC12913 CCRC14001	(262)	
GM-080	(257)	
CCRC16100	(264)	
KLB58		GTAGGCTGCAACTCGCCTACACGAAGTCGGAATCGCTAGTAATCGCGGATCAGCACGCCG
PB4	(539)	
F31		GTAGGCTGCAACTCGCCTACACGAAGTCGGAATCGCTAGTAATCGCGGATCAGCACGCCG
Consensus	(1321)	
00110011001	(/	
		1381 1440
CCRC12913	(251)	
CCRC14001	(262)	
GM-080		
CCRC16100	(264)	CGGTGAATACGTTCCCGGGCCTTGTACACACCGCCCGTCACACCATGAGAGTTTGTAACA
		- ΑΡΑΓΙΚΑ ΚΑΤΑΡΡΙΤΙΡΡΟΓΙΚΕΙ ΕΙΤΕΤΕΤΑΡΑΡΑΡΟΓΙΡΡΟΓΙΓΡΟΔΙΑΡΟΔΙΑΓΙΑΓΑΓΑΙΤΙΤΟΤΑΔΟΔ
KLB58		COUTOAATACOTTCCCOGCCTTGTACACACCGCCCGTCACACCATGAGAGTTTGTAGAC
PB4	(539)	
	(539)	CGGTGAATACGTTCCCGGGCCTTGTACACACCGCCCGTCACACCATGAGAGTTTGTAACA

FIG. 3 (continued)

		1441	1500	
CCRC12913	(251)			
CCRC14001	(262)			
GM-080	(257)			
CCRC16100	(264)			
KLB58	(1421)	$\tt CCCGAAGCCGGTGGCGTAACCCTTTTAGGGAGCGAGCTGTCTAAGGTGGGACAAATGATT$		
PB4	(539)			
F31	(1423)	CCCGAAGCCGGTGGCGTAACCCTTTTAGGGAGCGAGCT	GTCTAAGGTGGGACAAATGATT	
Consensus	(1441)			
		1501 1537		
CCRC12913	(251)			
CCRC14001	(262)			
GM-080	(257)			
CCRC16100	(264)			
KLB58	(1481)	AGGGTGAAGTCGTAACAAGGTAGCTAAAGGAGA		
PB4	(539)			
F31	(1483)	AGGGTGAAGTCGTAACAAGG-AGCTGTAGGAGAACTA		
Consensus	(1501)			

FIG. 3 (continued)

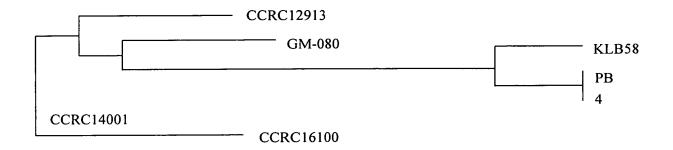


FIG. 4

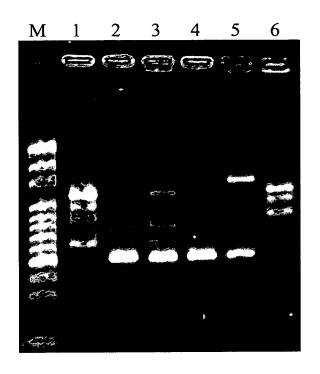


FIG. 5

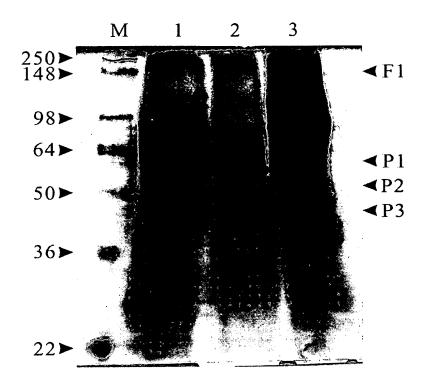


FIG. 6

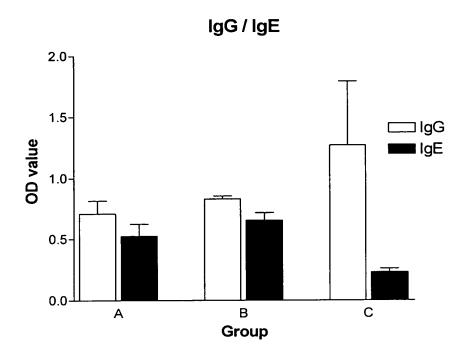


FIG. 7

BALF-Cell Count

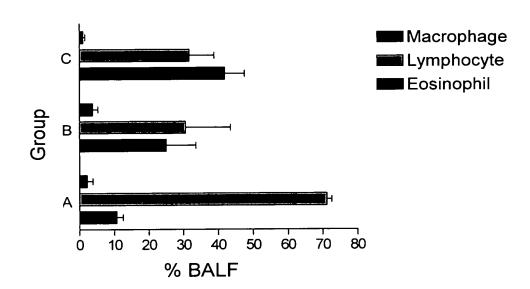


FIG. 8

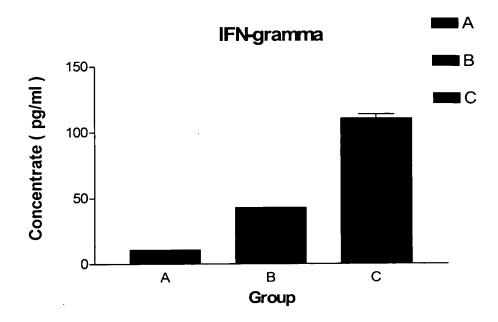


FIG. 9

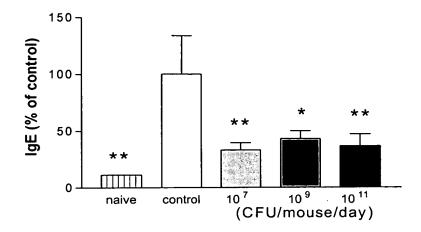


FIG. 10